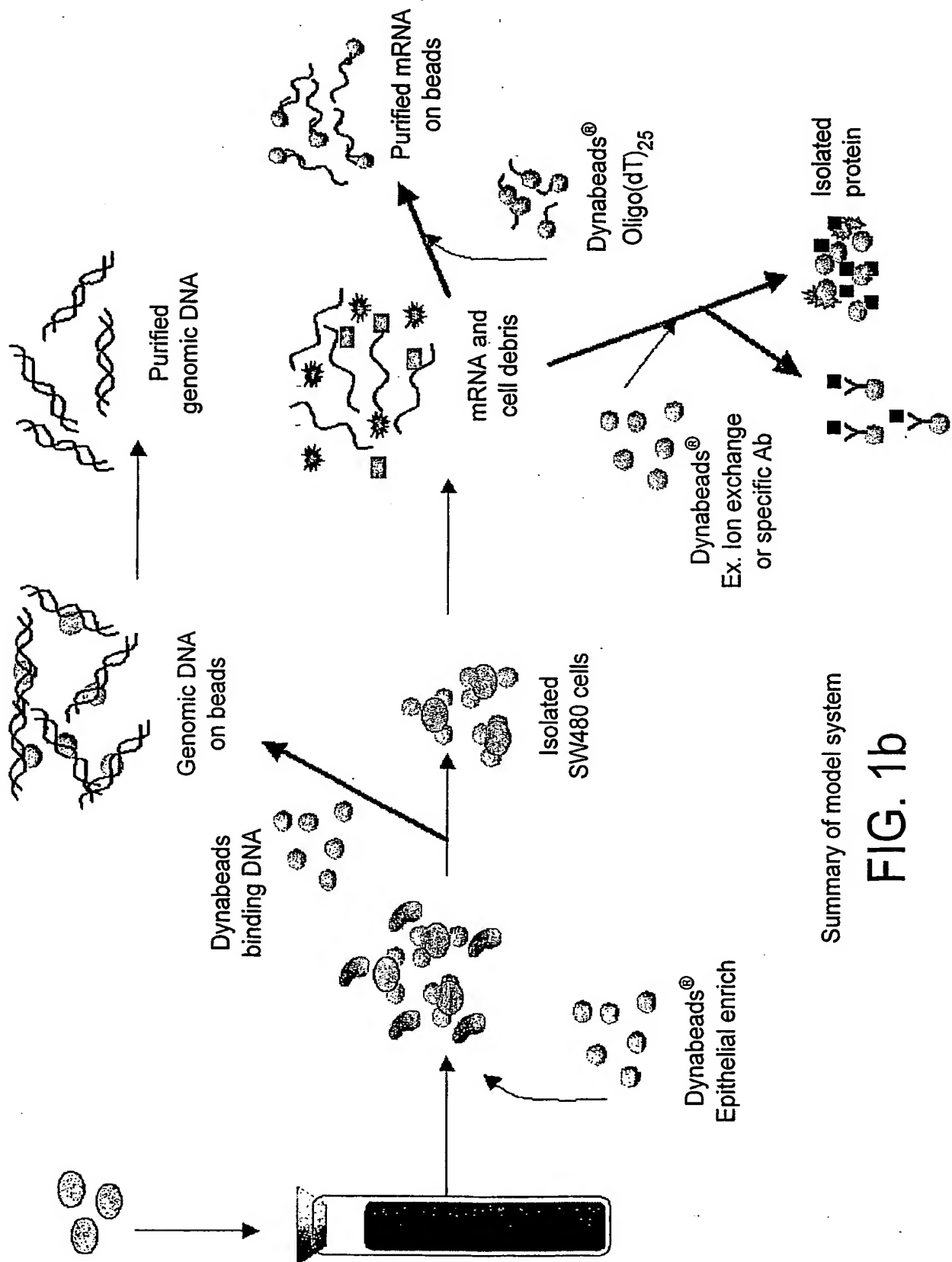


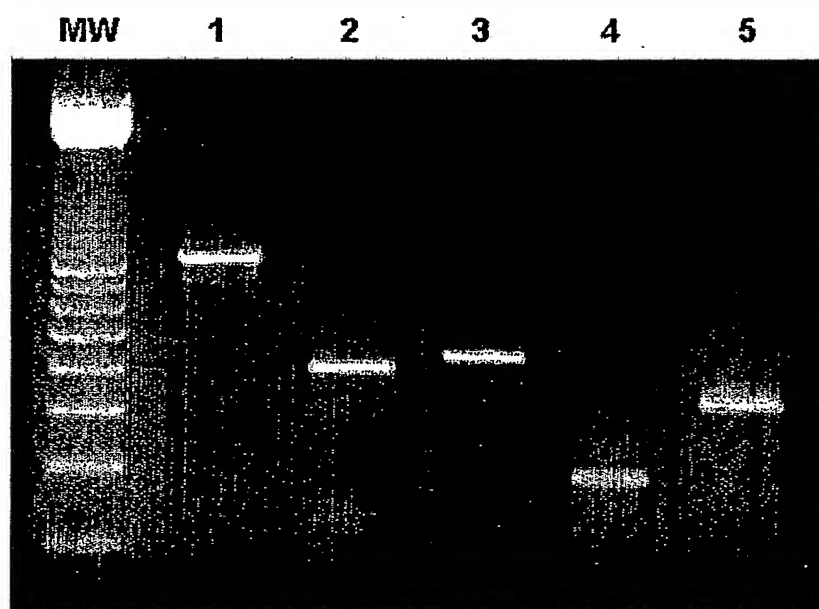


FIG. 1a



Summary of model system

FIG. 1b



Re-use of Solid-phase  
cDNA library in multiple  
PCR experiments,  
detecting different  
transcripts

Cultured Carcinoma Cells

| Lane | No. of cells | Amplicon | Round of PCR       |
|------|--------------|----------|--------------------|
| 1    | 10000        | ESX      | first (45 cycles)  |
| 2    | 10000        | Mucin1   | second (45 cycles) |
| 3    | 10000        | CK19     | third (30 cycles)  |
| 4    | 10           | CK19     | first (nested)     |
| 5    | 10           | ESX      | second (nested)    |

FIG. 2

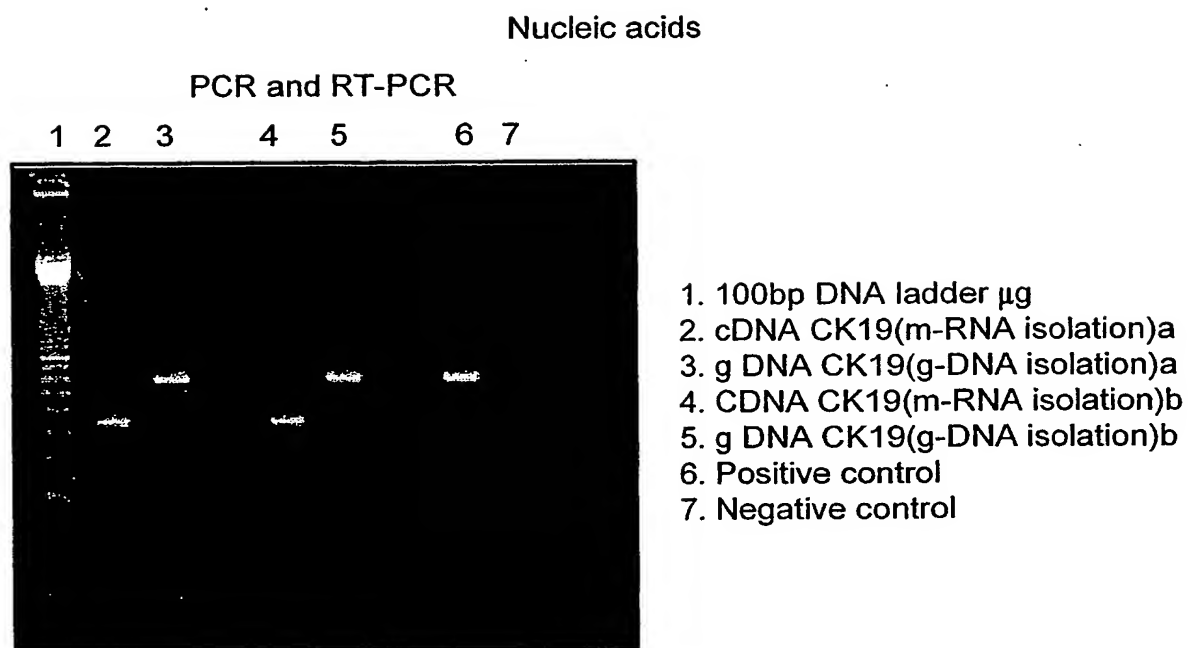


FIG. 3

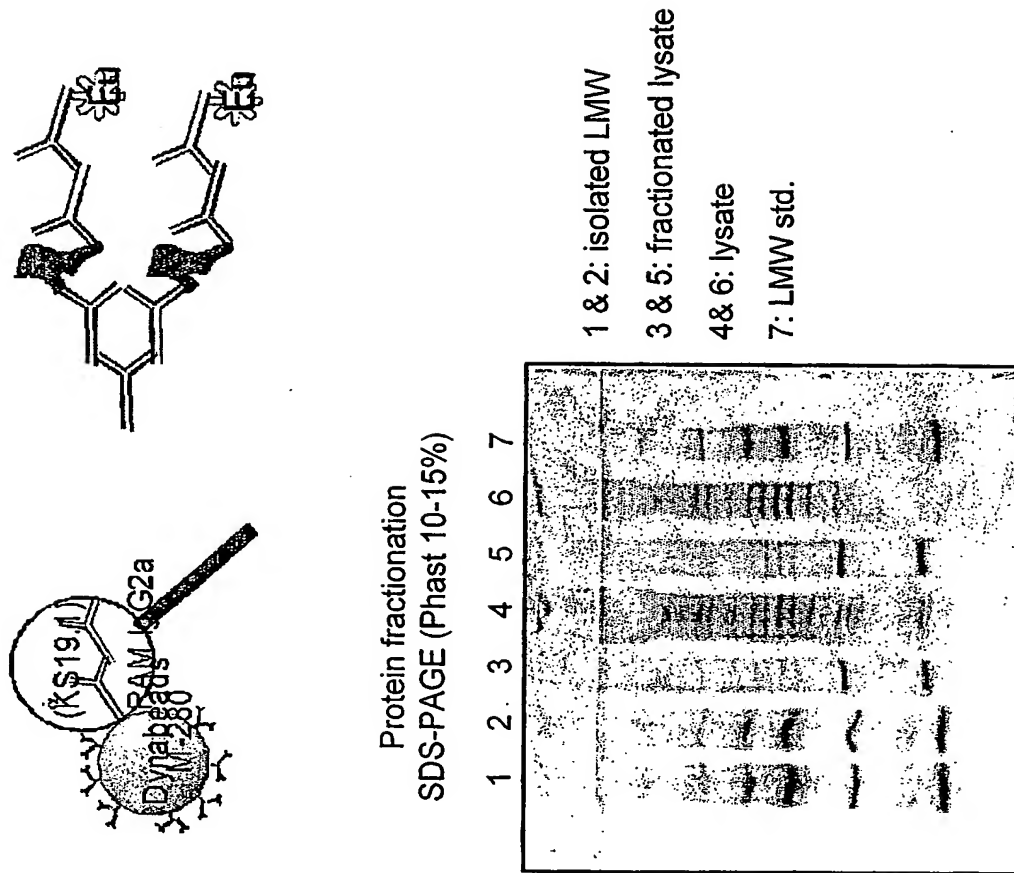
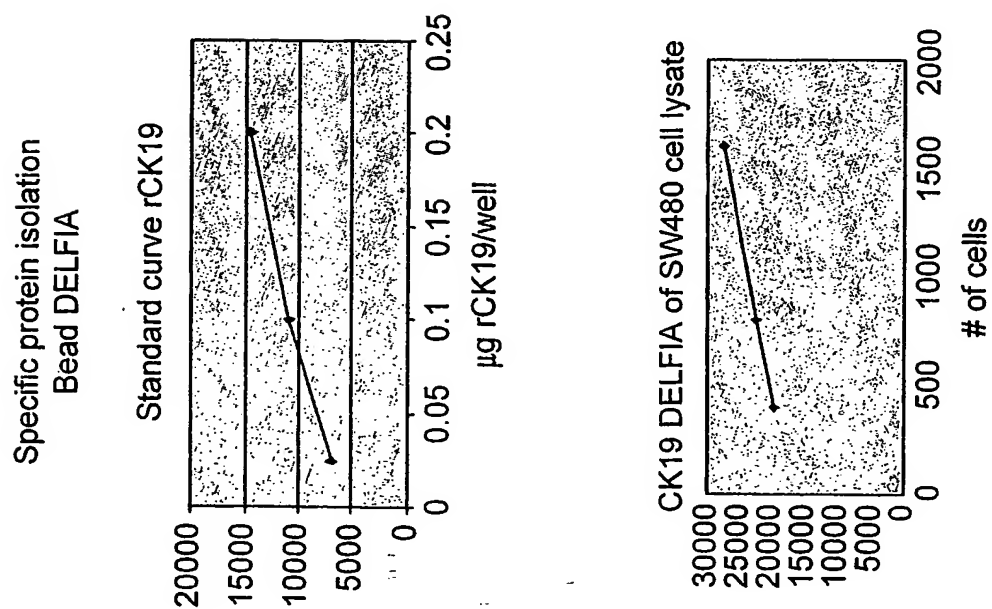


FIG. 4



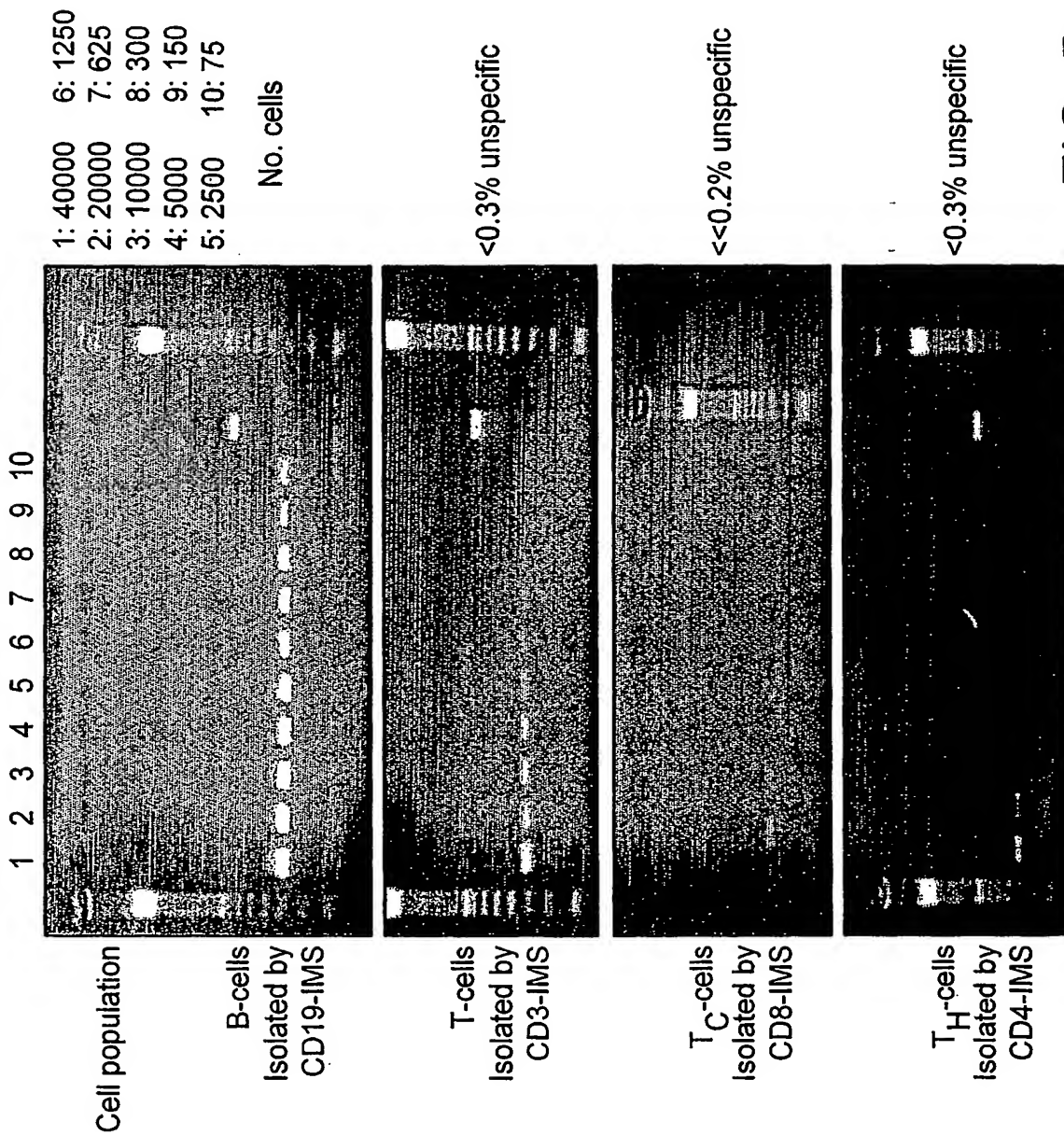


FIG. 5a

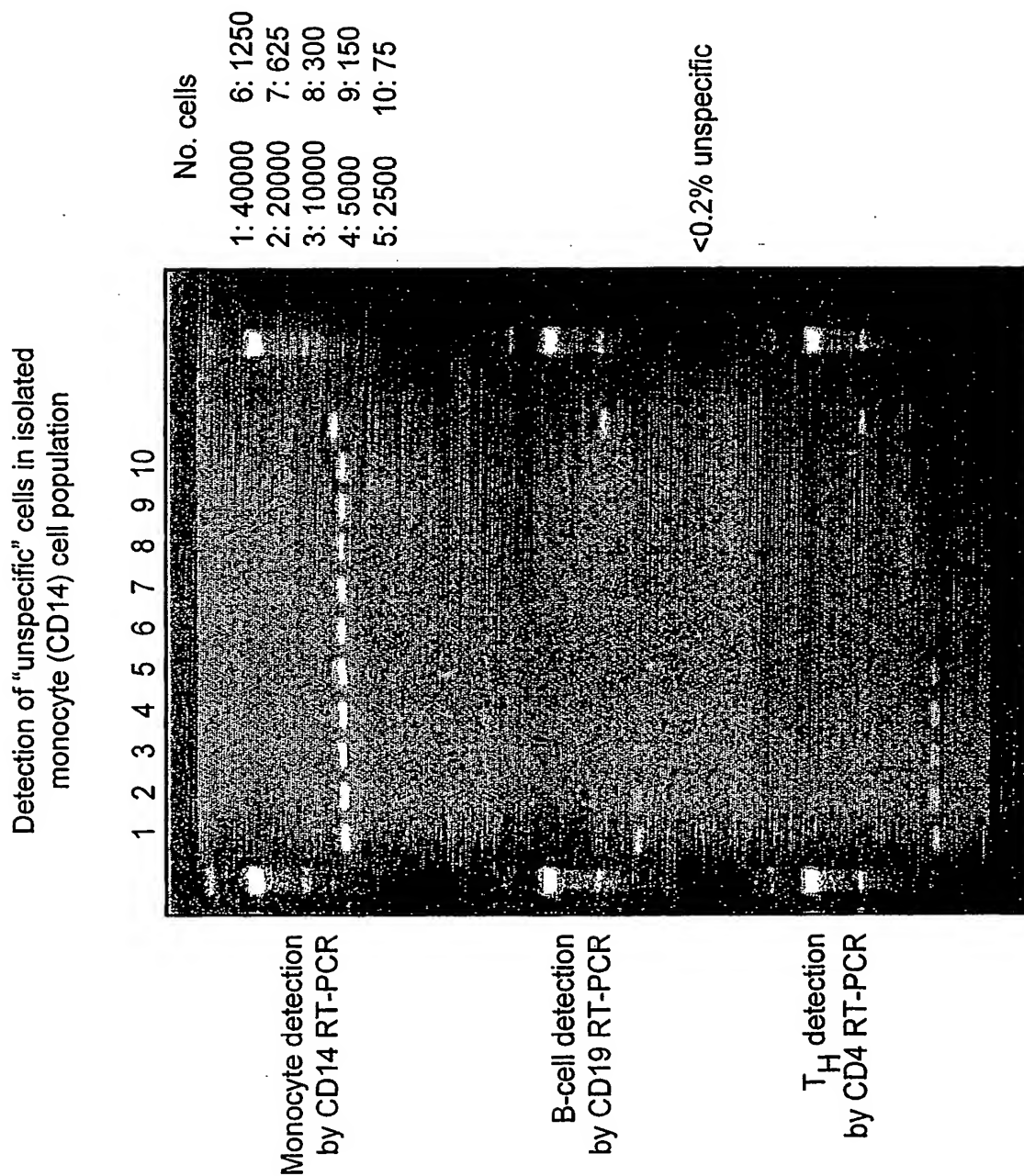
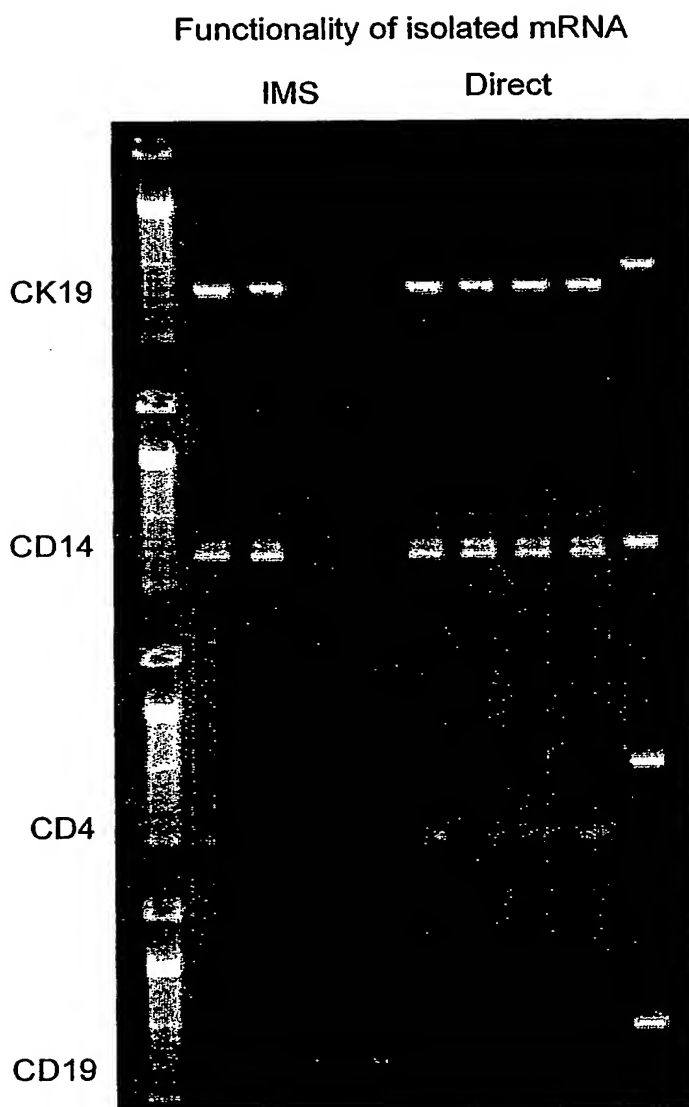


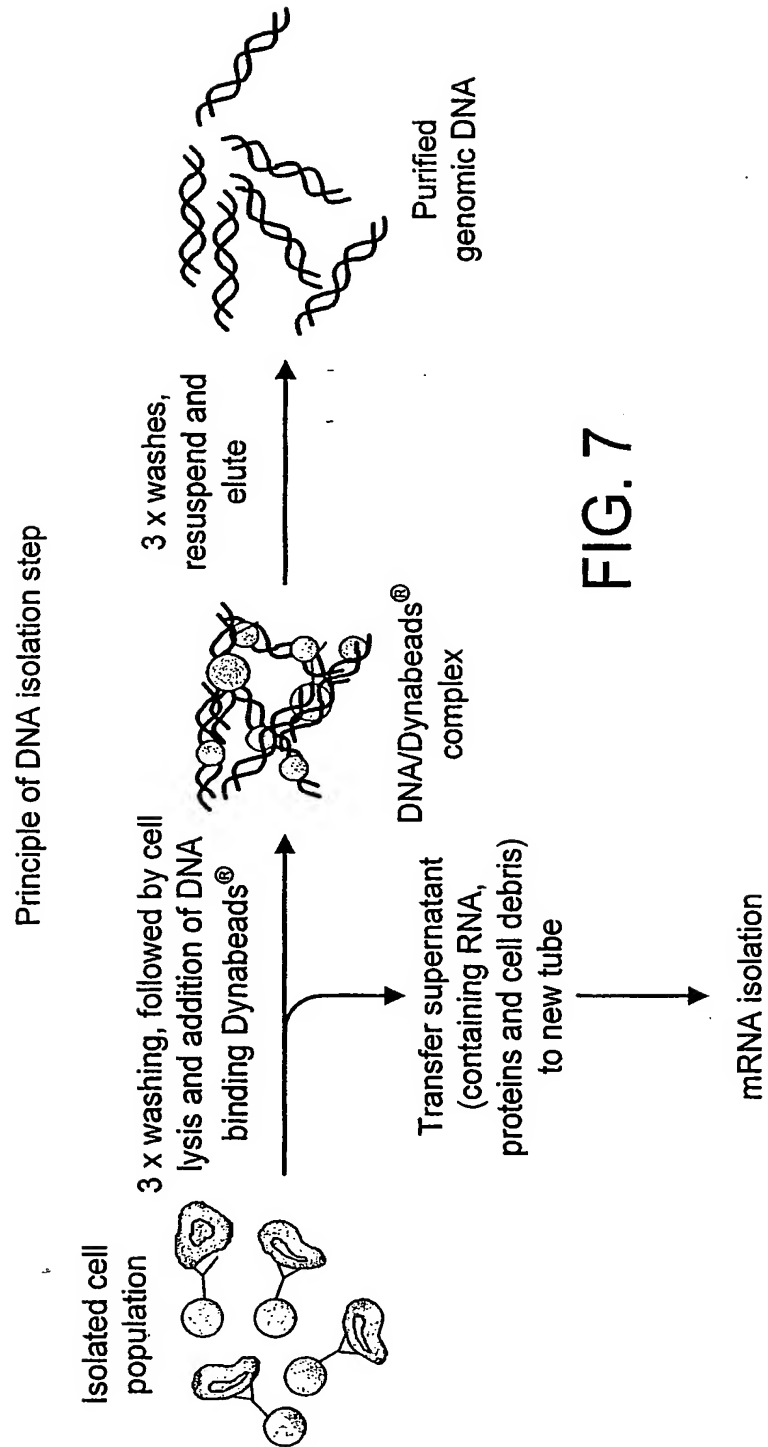
FIG. 5b

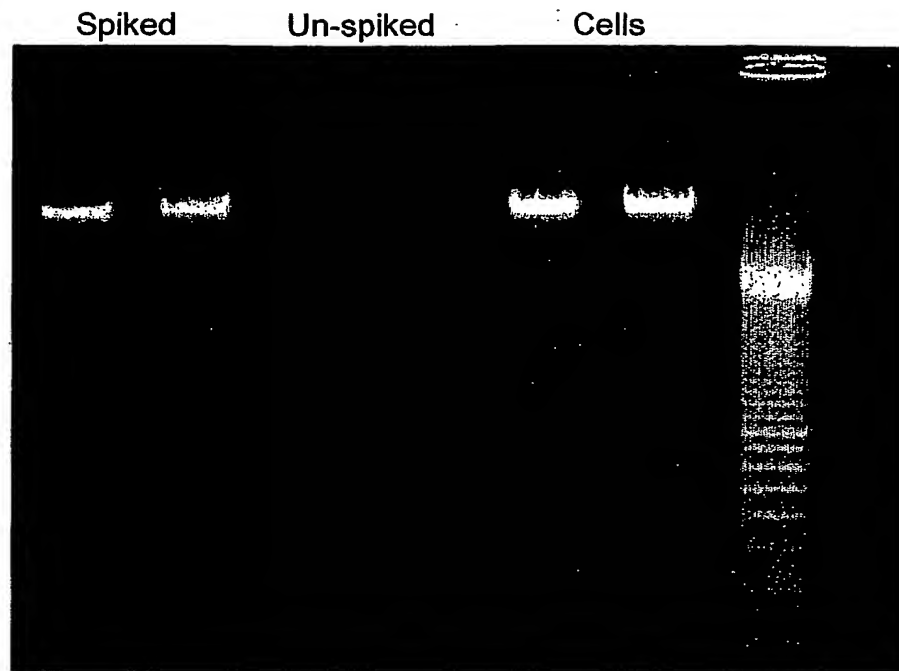


Template in RT-PCR:  
mRNA representing  
10000 carcinoma cells

FIG. 6







Integrity of isolated DNA:  
No degradation

FIG. 8

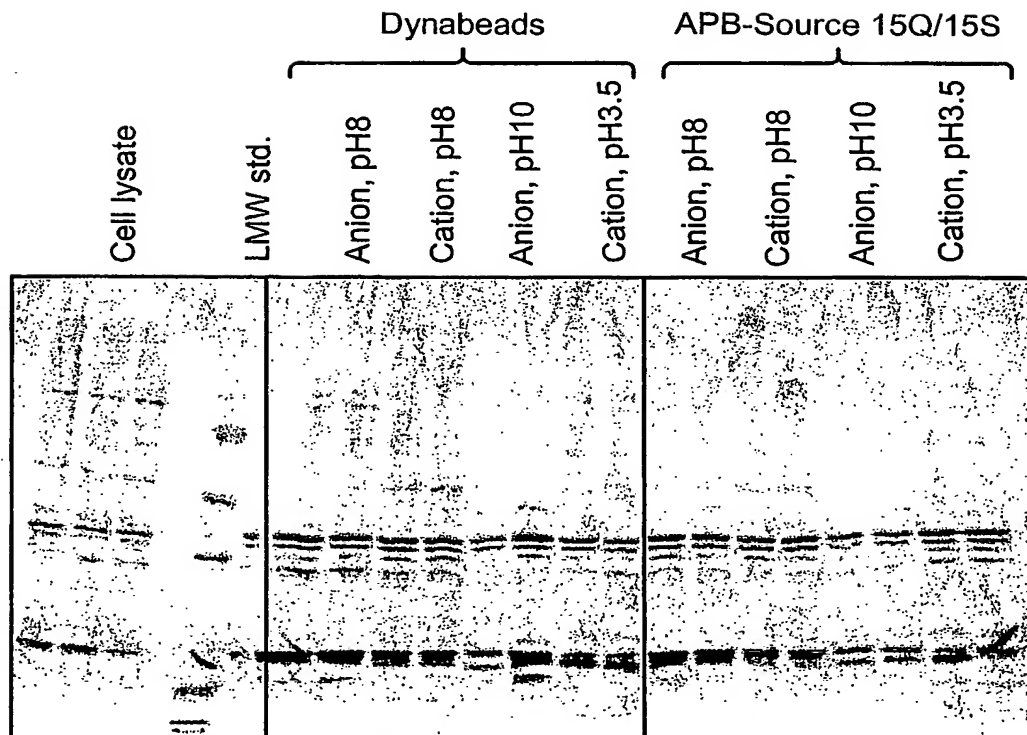


FIG. 9a

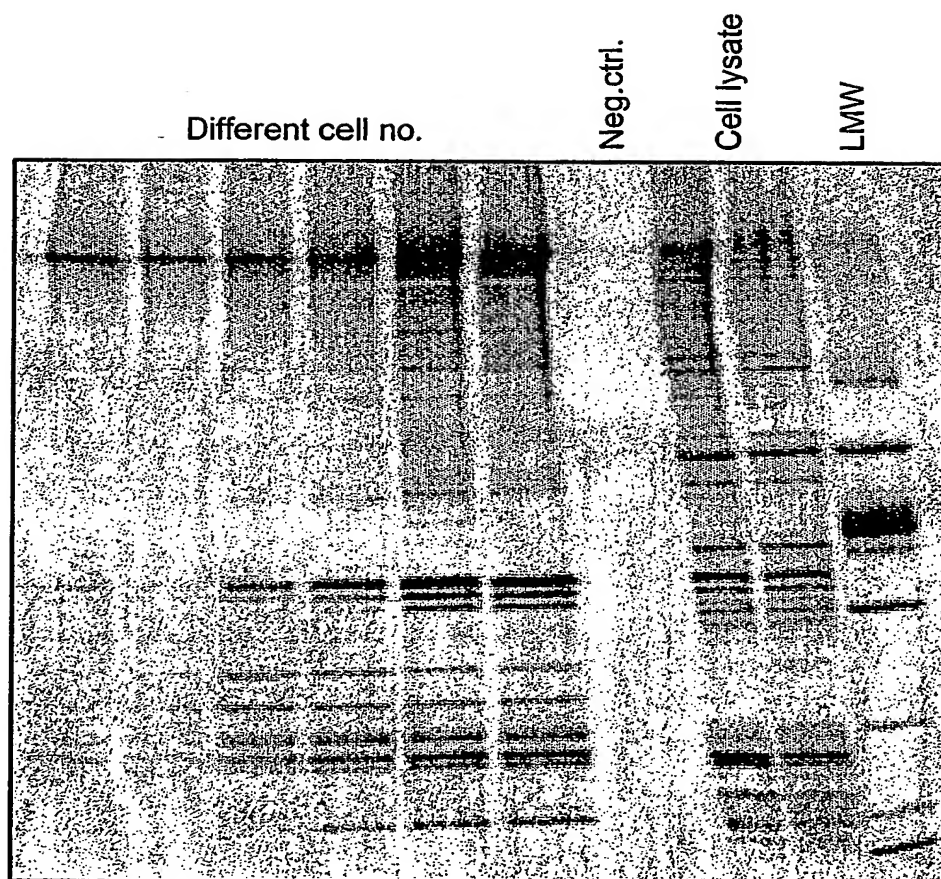


FIG. 9b

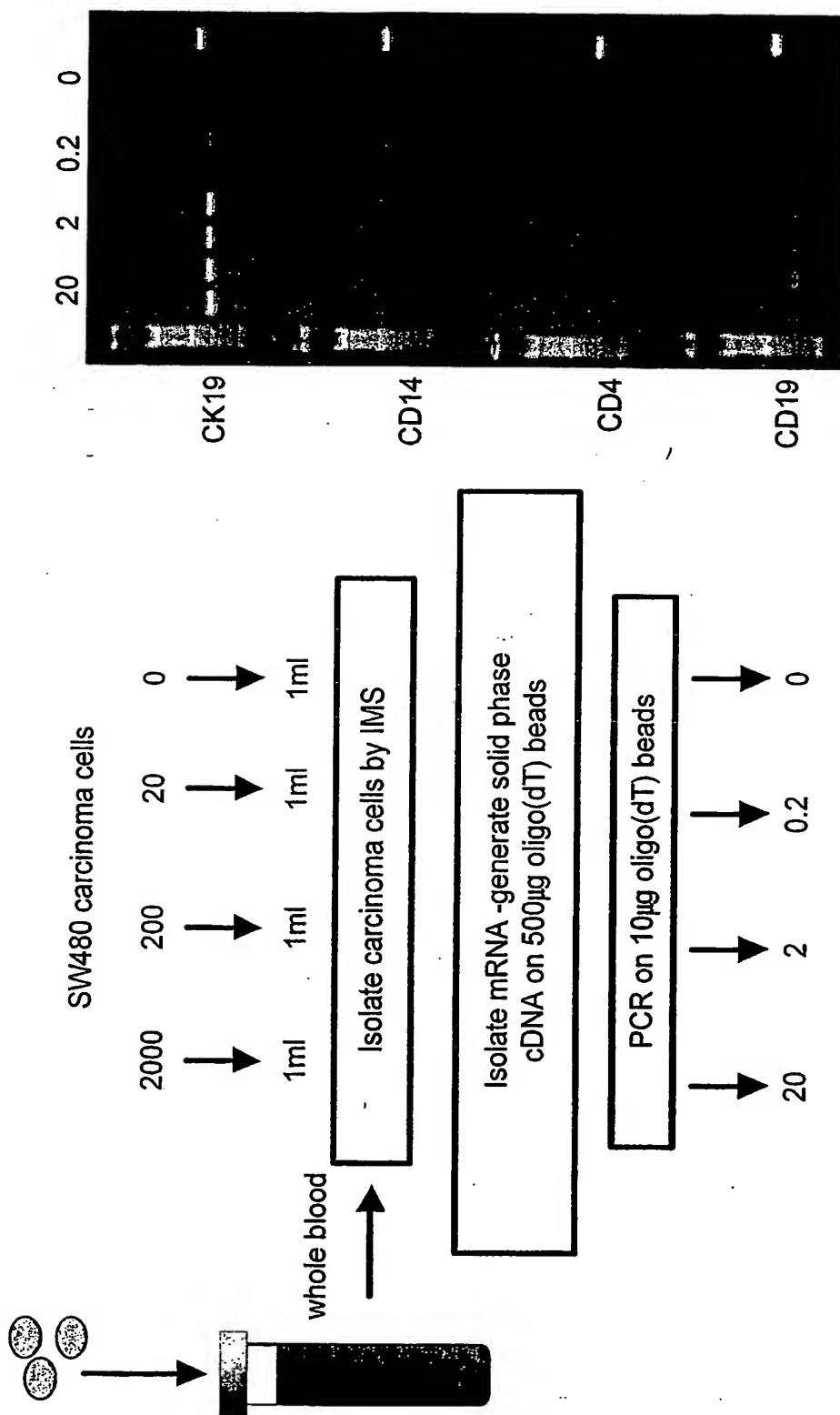


FIG. 10a

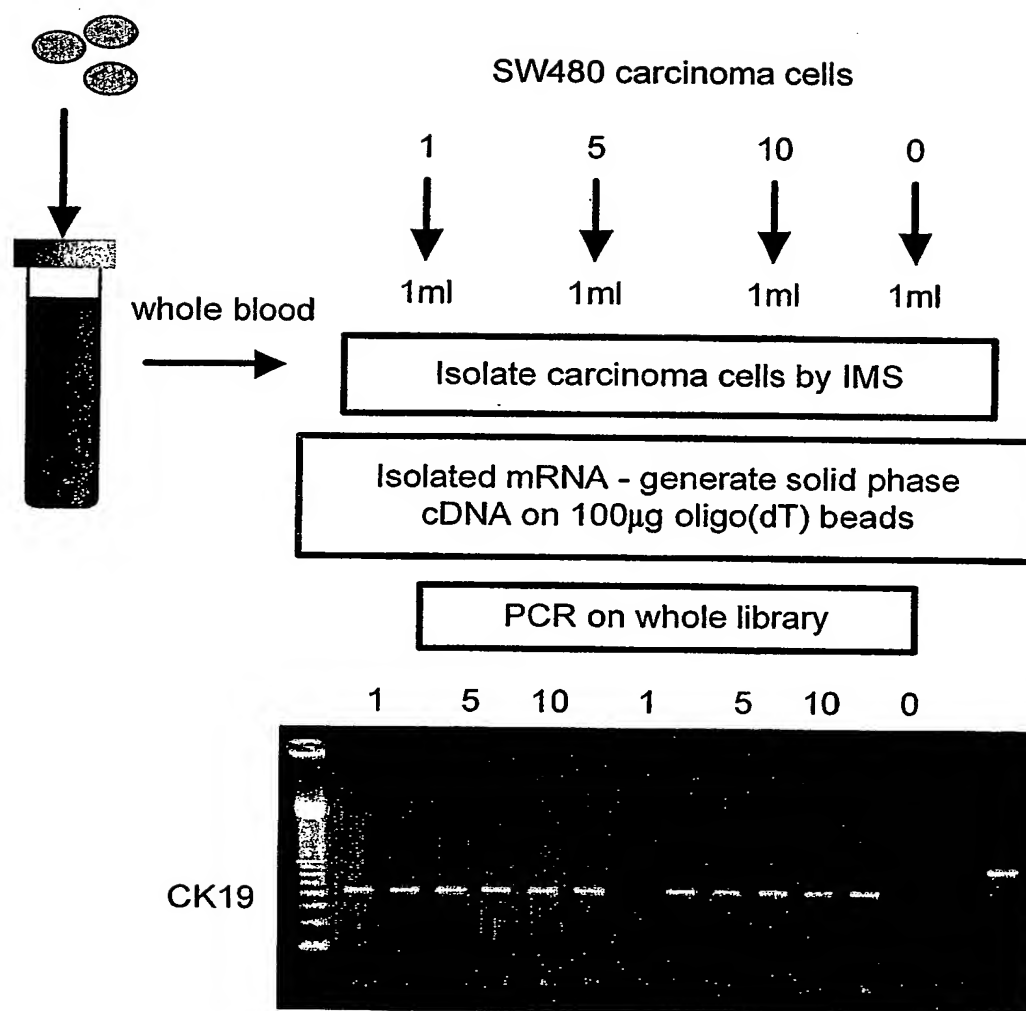


FIG. 10b

Flexible technology  
Re-Use of Solid Phase cDNA

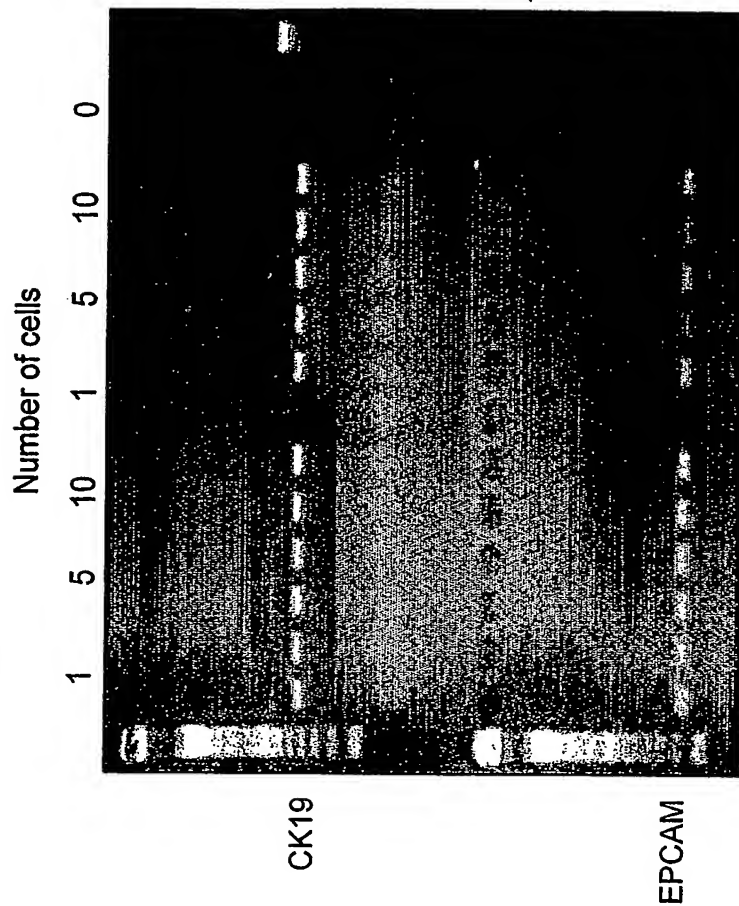
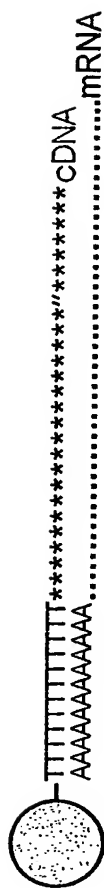


FIG. 11

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